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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=8; day=7; hr=9; min=50; sec=26; ms=60; ]

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Application No: 10534348 Version No: 1.1

Input Set:

Output Set:

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Finished: 2008-08-07 09:47:19.789

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Total Warnings: 0

Total Errors: 0

No. of SeqIDs Defined: 132

Actual SeqID Count: 132

# SEQUENCE LISTING

<110> SILENCE, Karen  
LAUWEREYS, Marc  
DE HAARD, Hans

<120> SINGLE DOMAIN ANTIBODIES DIRECTED AGAINST TUMOUR NECROSIS FACTOR-ALPHA AND USES THEREFOR

<130> A0848.70005US00

<140> 10/534,348

<141> 2005-05-09

<150> PCT/BE03/00192

<151> 2003-11-07

<150> US 60/425073

<151> 2002-11-08

<150> US 60/425063

<151> 2002-11-08

<150> EP 03447005.4

<151> 2003-01-10

<150> PCT/EP03/06581

<151> 2003-06-23

<150> PCT/EP03/07313

<151> 2003-07-08

<160> 132

<170> PatentIn version 3.1

<210> 1

<211> 115

<212> PRT

<213> Lama glama

<400> 1

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Thr Ser Gly Phe Asp Phe Ser Val Ser  
20 25 30

Trp Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ser Glu Ile Asn Thr Asn Gly Leu Ile Thr Lys Tyr Val Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr  
65 70 75 80

Leu Gln Met Asp Ser Leu Ile Pro Glu Asp Thr Ala Leu Tyr Tyr Cys  
85 90 95

Ala Arg Ser Pro Ser Gly Ser Phe Arg Gly Gln Gly Thr Gln Val Thr  
100 105 110

Val Ser Ser  
115

<210> 2

<211> 121

<212> PRT

<213> Lama glama

<400> 2

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Ser Ile Phe Arg Val Asn  
20 25 30

Ala Met Gly Trp Tyr Arg Gln Val Pro Gly Asn Gln Arg Glu Phe Val  
35 40 45

Ala Ile Ile Thr Ser Gly Asp Asn Leu Asn Tyr Ala Asp Ala Val Lys  
50 55 60

Gly Arg Phe Thr Ile Ser Thr Asp Asn Val Lys Lys Thr Val Tyr Leu

65		70		75		80
Gln Met Asn Val Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Asn						
	85		90		95	
Ala Ile Leu Gln Thr Ser Arg Trp Ser Ile Pro Ser Asn Tyr Trp Gly						
	100		105		110	
Gln Gly Thr Gln Val Thr Val Ser Ser						
	115		120			
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<212> PRT						
<213> Lama glama						
<400> 3						
Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly						
1	5		10		15	
Ser Leu Arg Leu Ser Cys Ala Thr Ser Gly Phe Thr Phe Ser Asp Tyr						
	20		25		30	
Trp Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val						
	35		40		45	
Ser Thr Val Asn Thr Asn Gly Leu Ile Thr Arg Tyr Ala Asp Ser Val						
	50		55		60	
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Tyr Thr Leu Tyr						
65		70		75		80
Leu Gln Met Asn Ser Leu Lys Ser Glu Asp Thr Ala Val Tyr Tyr Cys						
	85		90		95	
Thr Lys Val Val Pro Pro Tyr Ser Asp Asp Ser Arg Thr Asn Ala Asp						
	100		105		110	
Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser						
	115		120			

<210> 4

<211> 129

<212> PRT

<213> Lama glama

<400> 4

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Arg Thr Phe Ser Asp His  
20 25 30

Ser Gly Tyr Thr Tyr Thr Ile Gly Trp Phe Arg Gln Ala Pro Gly Lys  
35 40 45

Glu Arg Glu Phe Val Ala Arg Ile Tyr Trp Ser Ser Gly Asn Thr Tyr  
50 55 60

Tyr Ala Asp Ser Val Lys Gly Arg Phe Ala Ile Ser Arg Asp Ile Ala  
65 70 75 80

Lys Asn Thr Val Asp Leu Thr Met Asn Asn Leu Glu Pro Glu Asp Thr  
85 90 95

Ala Val Tyr Tyr Cys Ala Ala Arg Asp Gly Ile Pro Thr Ser Arg Ser  
100 105 110

Val Glu Ser Tyr Asn Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser  
115 120 125

Ser

<210> 5

<211> 127

<212> PRT

<213> Lama glama

<400> 5

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Arg Thr Phe Ser Ala His  
20 25 30

Ser Val Tyr Thr Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg  
35 40 45

Glu Phe Val Ala Arg Ile Tyr Trp Ser Ser Ala Asn Thr Tyr Tyr Ala  
50 55 60

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn  
65 70 75 80

Thr Val Asp Leu Leu Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val  
85 90 95

Tyr Tyr Cys Ala Ala Arg Asp Gly Ile Pro Thr Ser Arg Thr Val Gly  
100 105 110

Ser Tyr Asn Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120 125

<210> 6

<211> 124

<212> PRT

<213> Lama glama

<400> 6

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Ser Ile Phe Arg Val Asn  
20 25 30

Ala Met Gly Trp Tyr Arg Gln Val Pro Gly Asn Gln Arg Glu Phe Val  
35 40 45

Ala Ile Ile Thr Ser Ser Asp Thr Asn Asp Thr Thr Asn Tyr Ala Asp  
50 55 60

Ala Val Lys Gly Arg Phe Thr Ile Ser Thr Asp Asn Val Lys Lys Thr  
65 70 75 80

Val Tyr Leu Gln Met Asn Val Leu Lys Pro Glu Asp Thr Ala Val Tyr  
85 90 95

Tyr Cys Asn Ala Val Leu Gln Thr Ser Arg Trp Ser Ile Pro Ser Asn  
100 105 110

Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 7

<211> 123

<212> PRT

<213> Lama glama

<400> 7

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Thr Thr Ser Gly Arg Thr Ile Ser Val Tyr  
20 25 30

Ala Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
35 40 45

Ala Ser Ile Ser Gly Ser Gly Ala Ile Thr Pro Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Asn Pro Glu Asp Thr Ala Val Tyr Tyr Cys



85

90

95

Ala Ala Ser Arg Tyr Ala Arg Tyr Arg Asp Val His Ala Tyr Asp Tyr  
100 105 110

Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120

&lt;210&gt; 8

&lt;211&gt; 124

&lt;212&gt; PRT

&lt;213&gt; Lama glama

&lt;400&gt; 8

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Thr Arg Thr Phe Ser Arg Tyr  
20 25 30

Val Val Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
35 40 45

Ala Thr Ile Ser Trp Asn Gly Glu His Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Tyr Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr  
65 70 75 80

Leu Gln Met Gly Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Ala Arg Ser Phe Trp Gly Tyr Asn Val Glu Gln Arg Asp Phe Gly  
100 105 110

Ser Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser  
115 120

&lt;210&gt; 9

<211> 120

<212> PRT

<213> Lama glama

<400> 9

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Ser Ile Phe Arg Val Asn  
20 25 30

Ala Met Gly Trp Tyr Arg Gln Val Pro Gly Asn Gln Arg Glu Phe Val  
35 40 45

Ala Ile Ile Thr Asn Asp Thr Thr Asn Tyr Ala Asp Ala Val Lys Gly  
50 55 60

Arg Phe Thr Ile Ser Thr Asp Asn Val Lys Lys Thr Val Tyr Leu Gln  
65 70 75 80

Met Asn Val Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Asn Thr  
85 90 95

Val Leu Gln Thr Ser Arg Trp Asn Ile Pro Thr Asn Tyr Trp Gly Gln  
100 105 110

Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 10

<211> 120

<212> PRT

<213> Lama glama

<400> 10

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Ser Ile Phe Arg Val Asn  
20 25 30

Ala Met Gly Trp Tyr Arg Gln Val Pro Gly Asn Gln Arg Glu Phe Val  
35 40 45

Ala Ile Ile Ser Gly Asp Thr Thr Asn Tyr Ala Asp Ala Val Lys Gly  
50 55 60

Arg Phe Thr Ile Ser Thr Asp Asn Val Lys Lys Thr Val Tyr Leu Gln  
65 70 75 80

Met Asn Val Leu Glu Ser Glu Asp Thr Ala Val Tyr Tyr Cys Asn Ala  
85 90 95

Val Leu Gln Thr Ser Arg Trp Ser Ile Pro Ser Asn Tyr Trp Gly Gln  
100 105 110

Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 11

<211> 116

<212> PRT

<213> Lama glama

<400> 11

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ala Cys Val Ala Ser Gly Ser Ile Phe Ser Ile Asp  
20 25 30

Val Met Gly Trp Tyr Arg Gln Ala Pro Gly Gln Gln Arg Glu Leu Val  
35 40 45

Ala Thr Ile Thr Asn Ser Trp Thr Thr Asn Tyr Ala Asp Ser Val Lys  
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Val Val Tyr Leu  
65 70 75 80

Gln Met Asn Ser Leu Lys Leu Glu Asp Thr Ala Val Tyr Tyr Cys Asn  
85 90 95

Ala Arg Arg Trp Tyr Gln Pro Glu Ala Trp Gly Gln Gly Thr Gln Val  
100 105 110

Thr Val Ser Ser  
115

<210> 12

<211> 115

<212> PRT

<213> Lama glama

<400> 12

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Thr His  
20 25 30

Trp Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ser Thr Ile Asn Thr Asn Gly Leu Ile Thr Asp Tyr Ile His Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Lys Ser Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Leu Asn Gln Ala Gly Leu Ser Arg Gly Gln Gly Thr Gln Val Thr  
100 105 110

Val Ser Ser

&lt;210&gt; 13

&lt;211&gt; 126

&lt;212&gt; PRT

&lt;213&gt; Lama glama

&lt;400&gt; 13

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
 1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Arg Arg Thr Phe Ser Gly Tyr  
 20 25 30

Ala Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
 35 40 45

Ala Val Val Ser Gly Thr Gly Thr Ile Ala Tyr Tyr Ala Asp Ser Val  
 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Glu Asn Thr Val Tyr  
 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Gly Leu Tyr Tyr Cys  
 85 90 95

Ala Val Gly Pro Ser Ser Ser Arg Trp Tyr Tyr Arg Gly Ala Ser Leu  
 100 105 110

Val Asp Tyr Trp Gly Lys Gly Thr Leu Val Thr Val Ser Ser  
 115 120 125

&lt;210&gt; 14

&lt;211&gt; 123

&lt;212&gt; PRT

&lt;213&gt; Lama glama

<400> 14

Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Glu Phe Glu Asn His  
20 25 30

Trp Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ser Thr Val Asn Thr Asn Gly Leu Ile Thr Arg Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Tyr Thr Leu Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Lys Ser Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Thr Lys Val Leu Pro Pro Tyr Ser Asp Asp Ser Arg Thr Asn Ala Asp  
100 105 110

Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 15

<211> 124

<212> PRT

<213> Lama glama

<400> 15

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Gly Thr Leu Ser Ser Tyr  
20 25 30

Ile Thr Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
35 40 45

Gly Ala Val Ser Trp Ser Ser Ser Thr Ile Val Tyr Ala Asp Ser Val  
50 55 60

Glu Gly Arg Phe Thr Ile Ser Arg Asp Asn His Gln Asn Thr Val Tyr  
65 70 75 80

Leu Gln Met Asp Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Ala Arg Pro Tyr Gln Lys Tyr Asn Trp Ala Ser Ala Ser Tyr Asn  
100 105 110

Val Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser  
115 120

<210> 16

<211> 124

<212> PRT

<213> Lama glama

<400> 16

Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Ala Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Gly Thr Phe Ser Ser Ile  
20 25 30

Ile Met Ala Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val  
35 40 45

Gly Ala Val Ser Trp Ser Gly Gly Thr Thr Val Tyr Ala Asp Ser Val  
50 55 60

Leu Gly Ar